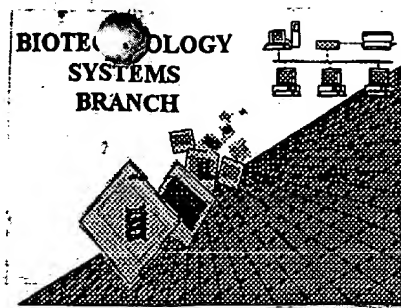


La Courne



#4
KAC
01-02-01

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

09/199,129

Source:

1635 RUSH

Date Processed by STIC:

1/4/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/199,129

DATE: 01/04/2001
TIME: 11:43:45

Input Set : D:\slreg.txt
Output Set: N:\CRF3\01042001\I199129.raw

Does Not Comply
Corrected Diskette Needed

ppr 1-5

1 <110> APPLICANT: Byrum, Joseph R.
2 La Rosa, Thomas J.
4 <120> TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
5 Plants
7 <130> FILE REFERENCE: 38-21(15075)B
9 <140> CURRENT APPLICATION NUMBER: US/09/199,129
9 <141> CURRENT FILING DATE: 1998-11-24
9 <160> NUMBER OF SEQ ID NOS: 5521

OK

global errors

ERRORED SEQUENCES

11 <210> SEQ ID NO: 1
12 <211> LENGTH: 254
E--> 13 <212> TYPE: nucleic acid *invalid response per new*
14 <213> ORGANISM: Glycine max *Sequence Rules (1.823)*
16 <223> OTHER INFORMATION: Clone ID: 700547901H1 *→ L2207 (see below)*
18 <400> SEQUENCE: 1
20 ggaaatctga agctacctta agcgaatcn anagggccaa nngcaacac tagtcacgac 60
22 ttcaagctac ttgattttc cttdnacte tgcactcttc attcgctgag attttcttcg 120
24 aagagagatt cgaagatgga tgaagagacc nnggggtgct ancgattgg aaaaggagtg 180
26 aaggaaaaagc ggttaccttc aagcacacag caggaaaatg gtcgtgggat cccgatgacn 240
28 aagggtcttca gaca *(see item 10 on error summary sheet)* 254
31 <210> SEQ ID NO: 2
32 <211> LENGTH: 285
E--> 33 <212> TYPE: nucleic acid
34 <213> ORGANISM: Glycine max *→ L2207*
36 <223> OTHER INFORMATION: Clone ID: 700547902H1
38 <400> SEQUENCE: 2
40 gggcgcttct ggngcgcgca cngetgttcc ttcgacgcgc ggcgngncgg gnggcgctgc 60
42 gccaaaccggg gactgcggcg ggtncctgt tctgcaacgg catcgngga agcccaaccn 120
44 ctagnctggc ggagttcacc ctgggcaacg ancatgantt ctacgaagtg agcctggtgg 180
46 acgggtacnn cctaccatc tccatnanch ccttcaaaag attccggaaa atycactacn 240
48 ccgggttgcy tgaacgagnt caacnccatg tgcccgcttg ggcct 285
51 <210> SEQ ID NO: 3
52 <211> LENGTH: 277
E--> 53 <212> TYPE: nucleic acid *→ L2207*
54 <213> ORGANISM: Glycine max
56 <223> OTHER INFORMATION: Clone ID: 700547903H1
58 <400> SEQUENCE: 3
60 cgggtgacca ttcattggaa gcttggaac cccagttaca tccagccctt tgatcgcatg 60
62 gtaactgtcc aacctgccc catacagaac cgcagtyagc ccactactaa gagggatcga 120
64 ggtgggctg gccacggat accttctggt gggcccatto gtgaaggccg ggcctcttag 180
66 gaacaccgag atcgccgggc aagcgggctc tctggccgcc ggtgggcttg tggatgcct 240
68 cagcctttgc ctcacaatct ntgggatttc atccttc 277
71 <210> SEQ ID NO: 4
72 <211> LENGTH: 219

check for n/s →
in sequence

check for n/s

use L2207 (header only - no response) WHENEVER
L2217, L2227, OR L2237 is shown. MANDATORY.

L2207-L2237
section.

RAW SEQUENCE LISTING

DATE: 01/04/2001

PATENT APPLICATION: US/09/199,129

TIME: 11:43:45

Input Set : D:\slreg..txt

Output Set: N:\CRF3\01042001\I199129.raw

```

E--> 73 <212> TYPE: nucleic acid
      74 <213> ORGANISM: Glycine max
      76 <223> OTHER INFORMATION: Clone ID: 700547904H1
      78 <400> SEQUENCE: 4
      80 agccatcacc atgggagcaq tgaattctccc agatcctcggc accgagattt tgattccggt 60
      82 ctgcgccacc attggaatag ggctcgcctt ctccagtggt gtcctcgtct ccaagggtta 120
      84 gctctccgct gccagagacg ctccccctaa cgcgcgcggc aaaaatgqct acaacgatta 180
      86 cctcaccgna gaagaggaag gcctcaacga tcacnacgt 219
      89 <210> SEQ ID NO: 5
      90 <211> LENGTH: 271

E--> 91 <212> TYPE: nucleic acid
      92 <213> ORGANISM: Glycine max
      94 <223> OTHER INFORMATION: Clone ID: 700547905H1
      96 <400> SEQUENCE: 5
      98 ctgcgcacc ctlatcctcg ttgtttctc ttlgcactc attcttggct ttttcgacga 60
      100 ctaccacct gcttcaaaag atgtacaaa ccggaagaag cactagtgtc atagtattca 120
      102 ttcttccagg ggataaaaata ttaggctctg caaaaatttc ttgggtatty gaatcccagg 180
      104 agatgccaaq gacttcttca atcaagtcga gtcataaggt ttctagagag caataggaaa 240
      106 gtgcttacta aaaaacaatct tatttaaatg c 271
      109 <210> SEQ ID NO: 6
      110 <211> LENGTH: 245

E--> 111 <212> TYPE: nucleic acid
      112 <213> ORGANISM: Glycine max
      114 <223> OTHER INFORMATION: Clone ID: 700547906H1
      116 <400> SEQUENCE: 6
      118 gcggcttcca atggcaccct tgcaattcac ctgcaccga ctcaaccaca ttgtncagg 60
      120 ctctctacc ttccacgcga accttctctc gaaaccgaag cctcatttgc aaagccactt 120
      122 tcttcttgtt ctgacgactt ctccaaaagn aggcgcgnt tcagcgcggn aancqccgna 180
      124 ctctcggnet cgtcccnca gttannngng cgtgncgaag angcgtttic ggagttggan 240
      126 anagt 245
      129 <210> SEQ ID NO: 7
      130 <211> LENGTH: 275

E--> 131 <212> TYPE: nucleic acid
      132 <213> ORGANISM: Glycine max
      134 <223> OTHER INFORMATION: Clone ID: 700547907H1
      136 <400> SEQUENCE: 7
      138 tcacaaaaca actttcaaac tctgagaaag aatggctgcc aacacattga tgagtactgc 60
      140 latctcagcc ttccacctct tctttcttct ctcaaaatcc agatttgcca ncgcagttcc 120
      142 tctttctagc ttgggtgtca ccaatgcctc ttcttctcgc ttctctatga gtgctgactg 180
      144 gatgccagcg nacctagac ctcccttanc tgatggttca gcacctgggt actttggatt 240
      146 cganccctct ngtcttgggt aagtaccaga gaatc 275
      149 <210> SEQ ID NO: 8
      150 <211> LENGTH: 273

E--> 151 <212> TYPE: nucleic acid
      152 <213> ORGANISM: Glycine max
      154 <223> OTHER INFORMATION: Clone ID: 700547909H1
      156 <400> SEQUENCE: 8
      158 anagcctagg gttcttttct tcttctgctc tcactccttg caaatctcgc gtaacgcatg 60
      160 cgttttctca ctcaatcgga gccacgact gatctatttg gaactcttgc aggtctgttg 120

```

same
error

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\slreg..txt
 Output Set: N:\CRF3\01042001\I199129.raw

```

162 ttttactgta ctcaatttga taaatraagg cccccccccn nnnnnnnnnn nnnntcccc 180
164 tgggtgttlat ggatgtgtcc atcgatgggg atcctgttga aaggalggtt tttagcttt 240
166 ctatgatgtt qctcccaaga ctgnagaaaa ctt 273
169 <210> SEQ ID NO: 9
170 <211> LENGTH: 235
E--> 171 <212> TYPE: nucleic acid
172 <213> ORGANISM: Glycine max
174 <223> OTHER INFORMATION: Clone ID: 700547910H1
176 <400> SEQUENCE: 9
178 agccattca aggataacga atttatgggt ggatcaattt caaacaggag gcgttctagg 60
180 caacgttcgg cctcaaatit tcttctcctg tatcaacaac ctcatctccc acaagccaa 120
182 gataatgggt ctgtggggca ttatgggtat tcattctcaga gctatagttg tggctgtgt 180
184 ccagaacaag taaagagctn ggataggaag tattaagggn taggtgatya ttata 235
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 265
E--> 189 <212> TYPE: nucleic acid
190 <213> ORGANISM: Glycine max
192 <223> OTHER INFORMATION: Clone ID: 700547911H1
194 <400> SEQUENCE: 10
196 caaaaaaccc aaatgcagql ttctccacta aaacttgggc qattttcana ttngtactt 60
198 tggcaccctt cactagatag ttctcaccat gtccgatact aaaagaagt ccaacaacag 120
200 cggttccaag cggagcctgc catcgtggac gaattcaagg qaaaatgaga gcgataacag 180
202 tgcaaaagaa ccaactttgg atggccaagg tgagaaatcc agtgacgctg agacacccca 240
204 caagagcaaa gtccaaaatg aaat 265
207 <210> SEQ ID NO: 11
208 <211> LENGTH: 273
E--> 209 <212> TYPE: nucleic acid
210 <213> ORGANISM: Glycine max
212 <223> OTHER INFORMATION: Clone ID: 700547913H1
214 <400> SEQUENCE: 11
216 gtcacaaccc aagcctccgc cgcgatttct cggccatgig cctcgaaatc gaggtttctg 60
218 accgctcttt ccggtaaaact caaccyggaa gtgactatga gqccaatggg gtgccctct 120
220 tctgctcttt tcaaggttga agccaagaag ggagagtggg tacctggctt ggcctcccca 180
222 acttacctca atggcactct tctgtgtgac aatggatttg accctctggg actagctgag 240
224 gaccagaga acttgaggtg gtacgttcaa gcc 273
227 <210> SEQ ID NO: 12
228 <211> LENGTH: 273
E--> 229 <212> TYPE: nucleic acid
230 <213> ORGANISM: Glycine max
232 <223> OTHER INFORMATION: Clone ID: 700547914H1
234 <400> SEQUENCE: 12
236 gcgtacaac aataaaaatc tccatttgt nttttcttct cttaacgcac accaacaata 60
238 cctctctctc atcggaatct ccaaaaagaa tggacaacaa aacgcagcag tccgagagta 120
240 agcaaaaacga caacgacgag gaagttgcgc caaaacgaca agaccctaac ccgtcgtctg 180
242 gcgggtgggg cttttcaccg ctctcgttct tctccgatct tcagaaggcc gccgctgttg 240
244 cagccgaaga gatctctcgc aatgctgctg tag 273
247 <210> SEQ ID NO: 13
248 <211> LENGTH: 277
E--> 249 <212> TYPE: nucleic acid

```

same

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\slreg..txt
 Output Set : N:\CRF3\01042001\I199129.raw

```

250 <213> ORGANISM: Glycine max
252 <223> OTHER INFORMATION: Clone ID: 700547916H1
254 <400> SEQUENCE: 13
256 agaacaagta gttgaqaact aagaaggaga agcaaatggc ttcctcaatg atctcttccc 60
258 cagctgttac cactgtcaac cgtgcccgtg ccggcatggt tgetccatc actggcctca 120
260 agtccatggc tggcttccc accagaaaga ccaacaatga cattanctcc attgctagca 180
262 acggtggaag agtgcnatgc atgcaqgtgt gggccaccag ttggcaagaa gaagtttgag 240
264 actcttccct acctgccaga ccttgatgat gcacatt 277
267 <210> SEQ ID NO: 14
268 <211> LENGTH: 275
E--> 269 <212> TYPE: nucleic acid
270 <213> ORGANISM: Glycine max
272 <223> OTHER INFORMATION: Clone ID: 700547917H1
274 <400> SEQUENCE: 14
276 qagcatttcn aatggagcaa gaaatttqgtg attcatttcc tttcgttggg nattgcattn 60
278 qcaatgatct ggattcattg attggcttgc nacgggnnct tgggtncctac tctqaacacn 120
280 aattttcaag atagctatgg gtgttgnata tttttntgtg aagaattaat gagcataatg 180
282 tttctlgaa agantggggg cagacgaalc tcattgtcaa ganttacaga gtttgagtgt 240
284 gtcaaaagca cttgtganga gtgttagcca gaagt 275
287 <210> SEQ ID NO: 15
288 <211> LENGTH: 273
E--> 289 <212> TYPE: nucleic acid
290 <213> ORGANISM: Glycine max
292 <223> OTHER INFORMATION: Clone ID: 700547920H1
294 <400> SEQUENCE: 15
296 attctctcca tatattatct caaacccctc tcacagaatg ggaagtgtg gaggaactga 60
298 ctatgttgca tacacttatg agaattctga gagagagcct tactggccat cagagaagct 120
300 taagatttcc atcactggng ctgggggttt tatcgcgtca cacatagctc ggcgcctcaa 180
302 gacagagggg cattacatta ttgcttctga ttggaagaaa aatgagcaca tgactyagga 240
304 catgttctgt gatgaattcc atcttgttga tct 273
307 <210> SEQ ID NO: 16
308 <211> LENGTH: 273
E--> 309 <212> TYPE: nucleic acid
310 <213> ORGANISM: Glycine max
312 <223> OTHER INFORMATION: Clone ID: 700547921H1
314 <400> SEQUENCE: 16
316 atcacaccat gccacgcta gtgacaatca agagaactcc ttcaagaagg gttctgctcg 60
318 ctctgccagt aaaagtcagg agaacaagtc atctggatta tcaaaagtcac cgactaatgc 120
320 aaataattat gggtcagttt ctctctcaag ntcaagtgtc cctgcaaaac gtactgagga 180
322 tgacatggat gattttgatc caagagggaac ttctaccaa acttcagctg gaaactctaa 240
324 cccaggttga tctcttttga caagattaat cgg 273
327 <210> SEQ ID NO: 17
328 <211> LENGTH: 245
E--> 329 <212> TYPE: nucleic acid
330 <213> ORGANISM: Glycine max
332 <223> OTHER INFORMATION: Clone ID: 700547924H1
334 <400> SEQUENCE: 17
336 gacacgtcca cggcgtcaaa ccaacttgag tgcgttccag gaaacacatc cctctccat 60
338 ttcatcact tctctgctct ctctttctcc caggaattcc caaacttatt ttacagcttc 120

```

Same

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\slreg..txt
 Output Set: N:\CRF3\01042001\I199129.raw

```

340 caatttccgt cagctaaacc atgcatcacc gcctcccaaa ttaggggttc cgacaggcgc 180
342 gtggagggaat cgtatggcgc cgtgaatccg cagcctctgc agttcgagga ccttgcctata 240
344 cccgt 245
347 <210> SEQ ID NO: 18
348 <211> LENGTH: 271
E--> 349 <212> TYPE: nucleic acid
350 <213> ORGANISM: Glycine max
352 <223> OTHER INFORMATION: Clone ID: 700547925H1
354 <400> SEQUENCE: 18
356 tttcttccca ttccaccct cgtcttccct tgtttccggt gagggccaac caaacaatat 60
358 gtttaacaag yaaccaacct caaaacataa catcctacgc caaacaacac ttgacctcc 120
360 ttcatatcta taccctnnnn nnnnnnnnnn ntctattttc aaattctaca tcatgggcac 180
362 qgaggttcta cggccacaag attgtttcac ccaacgcac ggtgttccac cgcctggctt 240
364 ttcccggcga agaacctatg gtaccacca c 271
367 <210> SEQ ID NO: 19
368 <211> LENGTH: 270
E--> 369 <212> TYPE: nucleic acid
370 <213> ORGANISM: Glycine max
372 <223> OTHER INFORMATION: Clone ID: 700547926H1
374 <400> SEQUENCE: 19
376 ccttattttc ttccctagt caaggagatt catctctgca gcagggtgtt ttgaaaaga 60
378 aaaaatgclg aaqatttgc gcaattgggc tggatatgtg ggggttccaa caatggcagt 120
380 cattgcactg aaatgccct caattgaagt ggtgtgtgtt gacatctcta aatccaggut 180
382 tgcagcatgg aacagtgaac agctccctat ctatgagcca ggccttgatg atgtggtgaa 240
384 qcaatgtcgt qgcaagaacc tttcttcag 270
387 <210> SEQ ID NO: 20
388 <211> LENGTH: 269
E--> 389 <212> TYPE: nucleic acid
390 <213> ORGANISM: Glycine max
392 <223> OTHER INFORMATION: Clone ID: 700547927H1
394 <400> SEQUENCE: 20
396 cnaqatttga aggttgcgta tgcctccat ctcccaactg tcttgcgcg tcttacatgc 60
398 aaatttctgt gaggaactgaa aactggcatt gttgggagaa cagggaagtgg taaatccact 120
400 ctcatacaaa cacttttccg aattgttgag cctactgccg gccaaqttat gattgacagc 180
402 atcaacatct ctccaatgga ctccatgatt tgaggtctag actaagcacc atcctccaga 240
404 ttccacaatg ttgaaggga cgtgagaa 269

```

same

FYI:

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

FYI:

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/199,129

DATE: 01/04/2001
TIME: 11:43:57

Input Set : D:\slreg..txt
Output Set: N:\CRF3\01042001\I199129.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:13 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:33 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:53 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:73 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:91 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:111 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:131 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:151 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:171 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:189 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:209 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:229 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:249 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:269 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:289 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:309 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:329 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:349 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:369 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:389 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:409 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:429 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:449 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:469 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:489 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:509 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:529 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:549 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:569 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:589 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:609 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:629 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:649 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:669 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:689 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:709 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:729 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:749 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:769 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:789 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:809 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:829 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:849 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:863 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:883 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:903 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/199,129

DATE: 01/04/2001
TIME: 11:43:57

Input Set : D:\slreg..txt
Output Set: N:\CRF3\01042001\I199129.raw

L:923 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:943 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:963 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:983 M:310 E: (3) Wrong or Missing Sequence Type, TYPE: